

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

CaltechDATA

RRID:SCR_017602

Type: Tool

Proper Citation

CaltechDATA (RRID:SCR_017602)

Resource Information

URL: <https://data.caltech.edu/>

Proper Citation: CaltechDATA (RRID:SCR_017602)

Description: Data and software repository from CalTech.

Resource Type: software repository, software resource, service resource, storage service resource, data repository

Keywords: Data, dataset, repository, CalTech

Availability: Restricted

Resource Name: CaltechDATA

Resource ID: SCR_017602

Alternate IDs: DOI:10.17616/R3sw99, r3d100012384, DOI:10.22002, DOI:10.25504/FAIRsharing.S09se7

Alternate URLs: <https://doi.org/10.17616/r3sw99>, <https://doi.org/10.22002/>, <https://dx.doi.org/10.22002/>, <https://fairsharing.org/10.25504/FAIRsharing.S09se7>, <https://doi.org/10.17616/R3SW99>

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260808T190111+0000

Ratings and Alerts

No rating or validation information has been found for CaltechDATA.

No alerts have been found for CaltechDATA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 11 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Chadly D, et al. (2026) Regenerative base editing enables deep lineage recording. bioRxiv : the preprint server for biology.

Ouadah Y, et al. (2026) Evolutionary basis of male same-sex sexual behavior by multiple pheromone switches in Drosophila. Current biology : CB, 36(7), 1716.

Fan C, et al. (2026) Bivalent mRNA booster encoding virus-like particles elicits potent polyclonal receptor-binding domain antibodies in pre-vaccinated mice. eLife, 14.

Kuintzle R, et al. (2025) Diversity in Notch ligand-receptor signaling interactions. eLife, 12.

Du R, et al. (2024) miRNA circuit modules for precise, tunable control of gene expression. bioRxiv : the preprint server for biology.

Xia S, et al. (2024) Synthetic protein circuits for programmable control of mammalian cell death. Cell, 187(11), 2785.

Briney KA, et al. (2024) Measuring data rot: An analysis of the continued availability of shared data from a Single University. PloS one, 19(6), e0304781.

Kuintzle R, et al. (2023) Diversity in Notch ligand-receptor signaling interactions. bioRxiv : the preprint server for biology.

da Veiga Beltrame E, et al. (2022) WormBase single-cell tools. Bioinformatics advances, 2(1), vbac018.

Segalin C, et al. (2021) The Mouse Action Recognition System (MARS) software pipeline for automated analysis of social behaviors in mice. eLife, 10.

B??guin JB, et al. (2020) Reduced volume and reflection for bright optical tweezers with radial Laguerre-Gauss beams. Proceedings of the National Academy of Sciences of the United States of America, 117(42), 26109.